# Sequence Listing.ST25 SEQUENCE LISTING IAD12 DA

IAP12 Rec'd PCT/PTO 09 MAY 2006

|                                  |                          |   | SEQUENC     | L L1311110 | IAP12 Rec    | d PCT/PTO  | $0.9 M\Delta$ |
|----------------------------------|--------------------------|---|-------------|------------|--------------|------------|---------------|
| <110>                            | Daw                      | Ventures, I<br>son, Elliot<br>ble, Kristi |             |            | 1711 12 1694 |            | U O INIT      |
| <120>                            | Met                      | hod and Sub                               | stances for | Diagnosing | Dyslexia     |            |               |
| <130>                            | 141                      | 60-1PCT                                   |             |            |              |            |               |
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| <210><br><211><br><212><br><213> | 1<br>3664<br>DNA<br>Homo | •   |             |            |              |            |               |
| <400><br>gaattaa                 | 1<br>agca                | ttttagcatt                                | ctttattaat  | ttttcaaagt | cactaggacc   | aaggataaca | 60            |
| attcato                          | catg                     | tgcatacaag                                | gccattctgt  | gtttcctact | cttgccttgg   | gctcatcatt | 120           |
| attaato                          | tgg                      | aattccattt                                | gttcttcact  | ttttgaatat | gtctgtttag   | ttgactgtag | 180           |
| tgccact                          | ggc                      | aggaccatgt                                | gcccaggaaa  | tccaagactc | atatttggac   | gaaagctatg | 240           |
| tccactt                          | ttc                      | aactagtacc                                | cctacccaaa  | ttaccatagc | aaccaaaaaa   | ttgcagatgc | 300           |
| ctacatt                          | cta                      | gaatcatgtt                                | ctaaagggat  | gtcatcattt | acaaaatgtc   | tttgttgagt | 360           |
| ctgaato                          | gtt                      | caaacaatag                                | caaaaaagga  | ttatttctct | cttggacatt   | tcaaagtact | 420           |
| atgacad                          | aaa                      | atatccaaga                                | cttgttatgg  | tgaggagcca | agtggaatgg   | aaaggacagc | 480           |
| tcatcco                          | ggc                      | ggctgggagt                                | gcatgcacac  | acatgccccc | tttttcttgc   | ctactaacag | 540           |
| gatctat                          | aga                      | aggcgtacat                                | aatgagtatg  | taggggactt | ggctgctttc   | agttaggaat | 600           |
| gagacac                          | tga                      | tatggttgga                                | atatagtaag  | agaaaaaggg | aggtctttct   | taaaaaacgg | 660           |
| ttttgtg                          | gtaa                     | aaatagagat                                | ggcacttaat  | ggatatcata | ttagcaggct   | ccctggacaa | 720           |
| atacata                          | agag                     | ccaaaacttc                                | tcatcgatta  | gccacctctt | caagtttagg   | ggttgaaaat | 780           |
| ctgaaac                          | caac                     | tacaaacatg                                | gtatctctct  | gaaaaggaga | taacgtaaaa   | gttatcacat | 840           |
| attaata                          | itaa                     | tgtgtatgaa                                | taaattgaca  | agctggttag | aaattagaaa   | taaaagtctt | 900           |
| gaggcaa                          | itaa                     | aagaggtaat                                | aacataggca  | aaaagagctc | ttcttctgga   | gagtggttgt | 960           |
| agatgga                          | igta                     | aacaagttta                                | ggtactgaac  | tgagaatagc | acatggatag   | accaattgtg | 1020          |
| gatgaag                          | ggag                     | actaaagaga                                | ggtttaacga  | atattgaaat | gaacctccag   | gtaggttgta | 1080          |
| tttatta                          | igtt                     | tgctgggaac                                | aagctgcttt  | tctctctct  | gtgaagcagg   | aaggcaaatt | 1140          |
| tctagtg                          | gct                      | ttccaaagga                                | aatgggaaat  | ctaaggaaat | ggtttgatac   | cagagtgttc | 1200          |

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Sequence Listing.ST25 tccttaggtt tattttaatg atggacttaa agatactttc ctatactcat gagctatgtt 1260 gtctctgata ttctttggta tattttacca aaaagataga ataggtgcca caagtattaa 1320 aaattttaga ctcctcagag cattacaaaa aacaagcaca aaatagaagc ctaatatgca 1380 gggaaagtca ctgaccatgc ccttggtact gctgattgta ttgcagagca agagatggac 1440 cctgagggta cttgaagcca acaagtttca cttctggaaa aagacttcag aatatgagtt 1500 taaaatataa aaagggaatt tgagccaaga cacaagaaca aactttttt gacaattata 1560 tctttattat tcctcttaca gagctacatt tactcttact aagtttcaga gtcaggtagt 1620 aatttacagt aagactgaat taccatccat aacgttagat gtccttattg aaacttcaac 1680 atcatttcca aatatcagca ttagcattgt gcttgacatt catttaacga agttactgaa 1740 aatctattaa gtataagaca tcagttattt ttaatagaag tttctgaaaa catttcagca 1800 aaatagcctg ttgagaaaaa tgtgtatgct gaaaaaaaaa aatgaacaaa taggaaagcc 1860 tggttcacaa acaggtgtca gggaaataga cagtactttt atagtaataa cataagaaca 1920 aacttettga aggtaagttt tattaaataa taggacaaca acaagataaa atgaettett 1980 cctgatattt atatattgat tgctggctgg tcataagact gtttttaggc aacgtgtttt 2040 gaaaaaccag aaagtctact accttgagtt ttcagccacg tgagaatagc aagattcagt 2100 gtttatactt gatagcatct taattaggcc tacaggcctc cctttcacat aactaccttc 2160 aagtttatga cagctcaaac tcacaattat cattatggag aagagagaag agttaagcta 2220 aaaacagacc actttcagag gacctgaaag caacgtaatc agtcacctat tgccatatac 2280 aagccacccc caaacataat gacttaaaac agcgatcatc tattattgct tatgagtctc 2340 tgagtcagct gaacattcct gctgatctgg gcttggttag gcttatttta gctgtgttca 2400 ttcttggtct gcagatagct gacaatcacc taggggctga ctgtaggcat tccagctgag 2460 atatgctctc tgtgtctttt atcctttagc aggaggaggc ttgctcacag ggtggttaca 2520. ggcatccaag agagtcagca taaatgtgaa aagtttccaa aatatcagat tcagtcctat 2580 gtaatctggt ttccattgca ttctcttggc cagagcaagt tgcaagacaa gtccaaattc 2640 aagaaggtca agaaatacac tccatctcca ggtaggagaa gctgcaaaga actgtgacaa 2700 tctatgacaa atagtatgtt caaagggaat aatatgggaa gatgtgccct ccgccaactt 2760 ctcagggaaa aatacagctt ttgtaatatt tagtaatata gactgtctaa tatttctaga 2820 gaaatctatg actttgagtt gaaatatctg aggccaacac tccaagcaat tttaaacaag 2880 tggtgacaga aattaccaga cacacatcaa gactcaagta taaagctata caatttaagg 2940 atgctcagca aatgttactg aattgactgg gtagtcccta aagagctgaa gaataaaaga 3000 tgttatgaga aatccaacaa taccaaatat aaattgcctc aggttctgaa atattcaata 3060 aagtattctc actgtagttc cttcagctta gctgatttgg actttggctg tgaaaacatt 3120

## Sequence Listing.ST25

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| atcctcagtg tttaaaaggt   | tggaaaattc 1 | tactgggtct | ttggcccaac | ctggaattaa | 3180 |
| atcctgatgc ttagaacctc a   | aagtctaaa a  | atcttctatt | gtcactttac | agagctattg | 3240 |
| aaacatatta ataaacttgt a   | atcatactga 1 | tttgattcta | atttttgtgg | gacattgttt | 3300 |
| aaaaattgtt gaaatgcata 1   | tatggaaaat t | tgatttttta | agtaaatgta | taacttttaa | 3360 |
| aattgtatcc tacatctaac 1   | ccaaataaa g  | ggtttaaaaa | caactatgag | caatataagt | 3420 |
| aatacattta aaatacattt a   | lagagaaaga t | taaggaaaaa | aggaatgact | catgaaggtt | 3480 |
| agtacacaat ctatgcatct 1   | gaatatttg c  | cacacttacc | aagtatttgg | ctccagggtt | 3540 |
| tctggcagct aatgcaaaga g   | jaggaacaga a | atcaagtttc | atggtattat | ctggtagact | 3600 |
| gtggaagcta tagcatttct g   | ccccctcat g  | gttttcacat | tccctttag  | agaacagcac | 3660 |
| aata  |              |            |            |            | 3664 |
| <210> 2<br><211> 22<br><212> DNA<br><213> Artificial<br><220><br><223> Artificial Seque | ence         |            | •          |            |      |
| <400> 2   | 10           |            |            |            | 2.2  |
| actaagaagt gcattagtcg g   | ) <b>g</b>   |            |            |            | . 22 |
| <210> 3<br><211> 20<br><212> DNA<br><213> Artificial                                    |              | ·          |            | •          | ·    |
| <220><br><223> Artificial Seque   | ence         |            |            |            |      |
| <400> 3<br>ttcctgtgct ctagcttgct  |              |            |            |            | 20   |
| <210> 4<br><211> 20<br><212> DNA<br><213> Artificial                                    |              |            | ·          |            |      |
| <220><br><223> Artificial Seque   | ence         |            | ,          |            |      |
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| <210> 5<br><211> 20<br><212> DNA<br><213> Artificial                                    |              |            |            |            |      |

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| 220                              | Se                              | quence Lis         | ting.ST25 |           |    |
|----------------------------------|---------------------------------|--------------------|-----------|-----------|----|
| <220><br><223>                   | Artificial Sequence             |                    |           |           |    |
| <400><br>ggttgc                  | 5<br>ctaa tcacgagaaa            |                    | -         |           | 20 |
| <210><br><211><br><212><br><213> | 6<br>25<br>DNA<br>Artificial    |                    | ,         |           |    |
| <220><br><223>                   | Artificial Sequence             |                    |           |           |    |
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| <210><br><211><br><212><br><213> | 7<br>25<br>DNA<br>Artificial    |                    |           |           |    |
| <220><br><223>                   | Artificial Sequence             | •                  |           |           |    |
| <400><br>ctagati                 | 7<br>tgaa ggccagaaaa catgc      |                    |           |           | 25 |
| <210><br><211><br><212><br><213> | 8<br>19<br>DNA<br>Artificial    |                    |           |           | •  |
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| <210><br><211><br><212><br><213> | 9<br>25<br>DNA<br>Artificial    |                    |           |           | ·  |
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| Met Va                           | l Arg Ser Gln Val Glu Trp L     | ys Gly Gln<br>Page |           | o Ala Ala |    |

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| 10 | Sequence<br>10 | Listing.ST25 |
|----|----------------|--------------|
|----|----------------|--------------|

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Gly Ser Ala Cys Thr His Met Pro Pro Phe Ser Cys Leu Leu Thr Gly 20 25 30

Ser Ile Glu Gly Val His Asn Glu Ala Ser Cys Lys Thr Ser Pro Asn  $\frac{35}{40}$ 

Ser Arg Arg Ser Arg Asn Thr Leu His Leu Gln Arg Asn Leu
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<400> 11

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Gly Ser Ala Cys Thr His Met Pro Pro Phe Ser Cys Leu Leu Thr Gly 20 25 30

Ser Ile Glu Gly Val His Asn Glu Ala Arg Asp Gly Pro 35 40 45

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|                 |                         | Sequence Listing.ST25                   |    |
|-----------------|-------------------------|---|----|
| <211><212><213> | 20<br>DNA<br>Artificial | • |    |
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| -9              | ,                       |   |    |
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| J.J J.          | 3 3 3 2 2 2 2 2         |   |    |
| <210>           | 19                      |   |    |
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| <220><br><223>                   | Artificial Sequence           |        |   |    |
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| <220><br><223>                   | Artificial Sequence           |        |   |    |
| <400><br>ctgact                  | 21<br>ctct tggatgcctg t       |        |   | 21 |
| <210><br><211><br><212><br><213> | 22<br>23<br>DNA<br>Artificial |        |   |    |
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| <210><211><211><212><213>        | 24<br>20<br>DNA<br>Artificial |        | - |    |
| <220><br><223>                   | Artificial Sequence           | Dama 7 |   |    |

| ٠, | W   | C | 119 | ណ    | 5/ | 049   | 70 | 2 |
|----|-----|---|-----|------|----|-------|----|---|
|    | ₩,₩ |   | -30 | ,0,0 | ~, | V 7.2 |    |   |
|    |     |   |     |      |    |       |    |   |

#### PCT/US2004/037850

#### Sequence Listing.ST25

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